



UNIVERSITI PUTRA MALAYSIA

***CROSSBREEDING BETWEEN CLEARFIELD® RICE
WITH WEEDY RICE UNDER VARIOUS CONDITIONS***

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By

ENGKU AHMAD KHAIRI BIN ENGKU ARIFF

**Thesis Submitted to the School of Graduate Studies, Universiti Putra Malaysia, in
Fulfillment of the Requirements for the Degree of Master of Science**

February 2016

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Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfilment of the requirement for the Degree of Master of Science.

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February 2016

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Rice is an important crop in our country as it is our staple food. Due to huge productivity losses because of weeds, the imidazolinone-resistant Clearfield® rice was developed to control it. Its close genetic relation with the weedy rice makes it a good candidate for hybridization, producing super weeds. The main objective of this study was to determine whether gene flow from Clearfield® rice to weedy rice can occur. This study has three experiments. In the first experiment, Clearfield® rice varieties (CL1, CL2) and weedy rice variants (V1, V2, V3, V4) were planted to observe the morphological characteristics. The second experiment was conducted in rice field for two seasons. The first (dry) season, two variants of weedy rice (V1, V2) and four variants (V1, V2, V3, V4) were used in and the second (rainy) season and were planted at a distance of 1m, 2m, 3m 4m and 5m from the Clearfield® rice. Seeds (F1) from weedy rice were collected and germinated in trays before were sprayed with OnDuty™ at day fourteen with a rate of 220 g/ha. The third experiment was determining hybrids using Simple Sequence Repeat (SSR) primer RM251 using leaves for the DNA extraction. The first experiment showed that weedy rice was morphologically superior to Clearfield® rice whereby it had double the number of tillers (more than 30) and almost 50 cm taller. In the second study after spraying OnDuty™, CL2 has significant difference at 20.38% compared to CL1 at 13.00% in second season. V1 showed the highest survival percentage, at 11.15% and 22.45% in both season. CL2 and V2 were the best combination of parent with 28.91% seedlings survived. About 80% seedlings survived from CL2V1 at the distance of 1m in second season. Higher number of overlapping period and wind speed in the second season were considered to affect survival percentages. The third study shows that molecular analysis has determined seven hybrids among the seedlings using primer RM251 with hybrids producing three bands. In conclusion, Clearfield® rice can hybridize with weedy rice under field condition and the percentages could increase with days of overlapping and wind speed.

Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan untuk Ijazah Sarjana Sains.

**KACUKAN ANTARA PADI CLEARFIELD®
DENGAN PADI ANGIN DI BAWAH PELBAGAI KEADAAN**

Oleh

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Padi merupakan tanaman penting di negara kita kerana ia adalah makanan ruji. Oleh kerana kerugian produktiviti yang banyak disebabkan rumpai, padi Clearfield® yang rintang imidazolinone telah dibangunkan untuk mengawalinya. Genetiknya yang berkait rapat dengan padi angin menjadikannya calon yang baik untuk penghibridan, menghasilkan rumpai super. Objektif utama kajian ini adalah untuk menentukan sama ada aliran gen daripada padi Clearfield® ke padi angin boleh berlaku. Kajian ini mempunyai tiga eksperimen. Dalam eksperimen pertama, varieti padi Clearfield® (CL1, CL2) dan varian padi angin (V1, V2, V3, V4) telah ditanam untuk melihat ciri-ciri morfologinya. Eksperimen kedua dijalankan di sawah untuk dua musim. Musim pertama (kering), dua varian (V1, V2) digunakan dan empat varian padi angin (V1, V2, V3, V4) telah digunakan dalam musim kedua (hujan) dan telah ditanam pada jarak 1m, 2m, 3m 4m dan 5m daripada padi Clearfield®. Benih (F1) dari padi angin telah dikumpulkan dan dicambahkan dalam dulang sebelum disemur dengan OnDuty™ pada hari empat belas dengan kadar 220 g / ha. Eksperimen ketiga adalah menentukan hibrid menggunakan primer Ulang Urutan Mudah (SSR) RM251 menggunakan daun untuk pengekstrakan DNA. Eksperimen pertama menunjukkan bahawa padi angin mempunyai morfologi lebih baik daripada padi Clearfield® di mana ia mempunyai dua kali ganda bilangan anak padi (lebih daripada 30) dan hampir 50 cm lebih tinggi. Dalam kajian kedua selepas menyemur OnDuty™, CL2 mempunyai perbezaan yang signifikan pada 20.38% berbanding dengan CL1 pada 13.00% pada musim kedua. V1 menunjukkan peratusan hidup yang paling tinggi, pada 11.15% dan 22.45% dalam kedua-dua musim. CL2 dan V2 adalah kombinasi terbaik induk padi dengan 28.91% benih terselamat. Kira-kira 80% benih terselamat dari CL2V1 pada jarak 1m dalam musim kedua. Tempoh bertindihan dan kelajuan angin yang lebih tinggi pada musim kedua telah dianggap mempengaruhi peratusan hidup. Kajian ketiga menunjukkan bahawa analisis molekul telah mengenalpasti tujuh kacukan daripada anak benih

menggunakan primer RM251 dengan kacukan menghasilkan tiga jalur. Kesimpulannya, padi Clearfield® boleh kacuk silang dengan padi angin di bawah keadaan lapangan dan kadar itu boleh meningkat disebabkan hari pertindihan dan kelajuan angin.



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This thesis was submitted to the Senate of Universiti Putra Malaysia and has been accepted as fulfilment of the requirement for the degree of Master of Science. The members of the Supervisory Committee were as follows:

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TABLE OF CONTENTS

	Page
ABSTRACT	i
ABSTRAK	ii
ACKNOWLEDGEMENTS	iv
APPROVAL	v
DECLARATION	vii
LIST OF TABLES	xii
LIST OF FIGURES	xiii
LIST OF ABBREVIATIONS	xv
 CHAPTER	
1 INTRODUCTION	1
2 LITERATURE REVIEW	3
2.1 Introduction of Rice	3
2.2 Production of Rice	3
2.3 Issues in Rice Production	4
2.4 Weedy rice	4
2.4.1 Origins of Weedy Rice	4
2.4.2 Characteristics of Weedy Rice	5
2.4.3 Problems Caused by Weedy Rice	5
2.5 Hybridization of weedy rice and commercial rice	5
2.6 Imidazolinone herbicides	6
2.7 Transgenic plants	7
2.7.1 Advantages of transgenic crops	8
2.7.2 Issues of transgenic crops	8
2.7.3 Hybridization of transgenic crops with wild relatives	9
2.8 Transgenic rice	9
3 MORPHOLOGICAL COMPARISON OF CLEARFIELD® RICE WITH WEEDY RICE	11
3.1 Introduction	11
3.2 Materials and Methods	12
3.2.1 Collection of rice seeds	12
3.2.2 Classification of weedy rice seeds	12
3.2.3 Vegetative and reproductive study	12
3.3 Results	13
3.3.1 Weedy rice classification	13
3.3.2 Seed morphology	13
3.3.3 Germination percentage	15
3.3.4 Height	15
3.3.5 Tillering ability	18
3.3.6 Reproductive study	20
3.4 Discussion	20

4	CONFIRMATION OF CROSSBREEDING BETWEEN WEEDY RICE AND CLEARFIELD® RICE WITH GENETIC MARKERS	22
4.1	Introduction	22
4.2	Materials and Methods	23
4.2.1	Field planting	23
4.2.2	Resistant study	25
4.2.3	Molecular analysis	26
4.2.3.1	DNA extraction	26
4.2.3.2	Primers for hybrid selection	26
4.2.3.3	PCR method	27
4.2.3.4	Optimization of SSR RM251	27
4.3	Results	28
4.3.1	Comparison of injury level between Clearfield® rice and weedy rice parents	28
4.3.2	Resistant study in first season	29
4.3.2.1	Conditions during the overlapping of panicle initiation in first season	29
4.3.2.2	Survivors of weedy rice F1 seedlings based on Clearfield® rice varieties	30
4.3.2.3	Survival rates of weedy rice F1 seedlings based on weedy rice variants	30
4.3.2.4	Survival rates of weedy rice F1 seedlings based on combination of Clearfield® Rice and weedy rice	31
4.3.3	Resistant study for second season	32
4.3.3.1	Conditions during the overlapping of panicle initiation in second season	32
4.3.3.2	Survivors of weedy rice F1 seedlings based on Clearfield® rice varieties	32
4.3.3.3	Survival rates of weedy rice F1 seedlings based on weedy rice variants	33
4.3.3.4	Survival rates of weedy rice F1 seedlings based on combination of Clearfield® Rice and weedy rice	34
4.3.4	Effects of distance toward survival rates of weedy rice seedlings	35
4.3.4.1	Effect on distance of survival rates of weedy rice F1 seedlings in first season	35
4.3.4.2	Effects of treatment combination between parents and distance on survival rates of weedy rice F1 seedlings in first season	36
4.3.4.3	Effect on distance of survival rates of weedy rice F1 seedlings in second season	37
4.3.4.4	Effects of treatment combination between parents and distance on survival rates of weedy rice F1 seedlings in second season	38
4.3.5	Confirmation of hybrids using molecular markers	39

4.3.5.1	DNA concentration and purity	39
4.3.5.2	Selection of suitable SSR primers for hybrid confirmation	40
4.3.5.3	Confirmation of Hybrids Using RM251	41
4.4	Discussion	41
5	SUMMARY, GENERAL CONCLUSION AND RECOMMENDATIONS FOR FUTURE RESEARCH	43
5.1	Summary and General Conclusion	43
5.2	Recommendation for Future Research	43
	REFERENCES	44
	APPENDICES	52
	BIODATA OF STUDENT	67
	LIST OF PUBLICATIONS	68

LIST OF TABLES

Table		Page
3.1	Weedy rice seed classification	13
3.2	Seed morphology of Clearfield® rice with weedy rice	15
3.3	Germination percentage	15
3.4	Mean height of Clearfield® rice and weedy rice variants from 7 DAT to 70 DAT	17
3.5	Mean number of tiller for Clearfield® rice with weedy rice variants from 7 DAT to 70 DAT	19
4.1	Forward and reverse primer sequences	27
4.2	Concentration and purity of Clearfield® rice and weedy rice parents	39

LIST OF FIGURES

Figure		Page
2.1	Imidazolinone herbicide family; a: Imazapyr b: imazapic c: imazethapyr d: imazamox e: imazaquin	7
3.1	Seeds of weedy rice and Clearfield® rice using QuickPHOTO MICRO 2.3	14
3.2	Height of Clearfield® rice and weedy rice variants from 7 DAT until 70 DAT	17
3.3	Number of tillers of Clearfield® rice and weedy rice variants from 7 DAT until 70 DAT	19
3.4	Panicle initiation of Clearfield® and weedy rice	20
4.1	Planting area for first season. R: Replicate. Rice type: CL1, CL2, V1 and V2	24
4.2	Planting area for second season. R: Replicate. Rice type: CL1, CL2, V1, V2, V3 and V4	25
4.3	Parents of Clearfield® and weedy rice at 1 week after spraying	29
4.4	Survival percentage of weedy rice F1 seedlings from different Clearfield® rice plots 1 WAT with OnDuty™	30
4.5	Survival percentage of weedy rice F1 seedlings from different variants 1 WAT with OnDuty™	31
4.6	Survival percentage of weedy rice F1 seedlings from different combination treatment of parents 1 WAT with OnDuty™	32
4.7	Survival percentage of weedy rice F1 seedlings from different Clearfield® rice plots 1 WAT with OnDuty™	33
4.8	Survival percentage of weedy rice F1 seedlings from different variants 1 WAT with OnDuty™	34
4.9	Survival percentage of weedy rice F1 seedlings from different combination treatment of parents 1 WAT with OnDuty™	35
4.10	Survival percentage of weedy rice seedlings based on planting distance 1 WAT with OnDuty™	36

4.11	Survival percentage of weedy rice based on treatment combination of parents and distance after 1 WAT with OnDuty™	37
4.12	Survival percentage of weedy rice F1 seedlings based on planting distance 1 WAT with OnDuty™	38
4.13	Survival percentage of weedy rice based on treatment combination of parents and distance after 1 WAT with OnDuty™	39
4.14	Amplified products from genomic DNA of Clearfield® rice and weedy rice parents.	40
4.15	Confirmation of hybrids. L1: Clearfield® rice. L2 to L3: Weedy rice. L4 to L7: Detected hybrids	41

LIST OF ABBREVIATIONS

®	Registered trademark
™	Trademark
μ	micro
ALS	acetolactate synthase
ANOVA	Analysis of variance
Bt	<i>Bacillus thuringiensis</i>
DAS	Day after seeding
DAT	Day after transplanting
DNA	Deoxyribonucleic acid
FAO	Food and Agriculture Organization
g	Gram
ha	Hectare
ht	Height
IGMORIS	Indian GMO research Information system
IMI	Imidazolinone
IRRI	International Rice Research Institute
ISAAA	International service for the Acquisition of Agri-biotech Application
L	Litre
LKPP	Lembaga Kemajuan Perusahaan Pertanian
m	metre
MARDI	Malaysian Agriculture Research and Development Institute
min	Minute
mm	millimeter
MRL	Maximum Residue Limit

mt	Million tons
PCR	Polymerase chain reaction
RAPD	Random amplified polymorphic DNA
Sdn Bhd	Sendirian Berhad
SSR	Simple sequence primer
SSLP	simple sequence length polymorphism
TAE	Tris/acetate/ Ethylenediaminetetraacetic acid
TBE	Tris/Borate/Ethylenediaminetetraacetic acid
Ti	Tillering ability
USD	United States Dollar

CHAPTER 1

INTRODUCTION

Rice is a very important crop in Asia as 90% of all rice production are consumed in this region (Gealy *et al.*, 2003). It is the main staple food in Malaysia and the third largest crop production after palm oil and rubber. However rice is very weak in term of competitiveness as yield loss caused by weeds can go up to 35% (Karim *et al.*, 2004). Under field conditions, weedy rice can absorb up to 60% of nitrogen (N) fertilizer that was applied (Burgos *et al.*, 2006). Most of the weeds that can cause serious economic problems are the wild *Oryza*. They compete for sun, nutrients and water with commercial rice (Chin *et al.*, 2007). Weedy rice is genetically related to cultivated rice (Gealy *et al.*, 2003). Because of the genetic similarity they shared, controlling weedy rice is very difficult. Most weedy rice have the same common trait such as taller plants, fewer tillers, easy shattering of the seeds, and earlier time of flowering (Chin *et al.*, 2007). The ease of shattering and seed dormancy of weedy rice can effect seed bank (Burgos *et al.* 2011) and increase managerial problem in the years to come. Compared to weedy rice, cultivated rice is less efficient in terms of N absorption efficiency while weedy rice can produce more biomass per one unit of N absorbed (Burgos *et al.*, 2006). During harvesting, grain taken from weedy rice can reduce the quality of milled rice due to the extra milling done to remove the red pigments from red rice seeds (Shivrain *et al.*, 2008).

The Clearfield® rice was developed specifically to control weeds in the rice fields. It is a type of naturally genetically modified rice that is resistant to imidazolinone based herbicides (Croughan, 2003). In Malaysia two varieties were released, MR220-CL1 and MR220-CL2. Although this technique is effective in controlling weedy rice it also has drawbacks. As Clearfield® rice and weedy rice are genetically related, these rice have the possibility to hybridize. Natural hybridization can occur depending on the factors such as genetic and environments. In terms of genetics, weedy rice and common cultivated rice are distinctly related. Although rice is self-pollinated, rice pollen can travel long distances from their mother plant. This can cause gene flow to occur. Gene flow occurrences are very low, less than 1% or less than 200 plants per hectare but the statistics can change due to type of weedy rice and cultivated rice that are within the vicinity of the area (Shivrain *et al.*, 2008).

The main concern of this hybridization is the production of progenies that have the same resistance as Clearfield® rice and thus will cause a problem for rice production. Countries like Brazil (Roso *et al.*, 2010), Greece (Kaloumenos *et al.*, 2013), and United State (Shivrain *et al.*, 2008) all have reported gene flow from Clearfield® rice to weedy rice under field condition. For our country that also uses Clearfield® rice, the chances of gene flow are almost certain. There are three objectives for this study. The objectives of this study are:

1. To determine the morphology differences of common weedy rice with Clearfield® rice

2. To study the possibility of hybridization between Clearfield® rice and weedy rice in field condition using genetic markers as confirmation.
3. To determine the distance factor that increases the hybridization frequency between Clearfield® rice with weedy rice



REFERENCES

- Ahmed, Q. N., Roy, R., Kamaruzaman, M. & Othman, A. S. (2012). Vegetative and Reproductive Growth of Weedy Rice in Selangor, Malaysia: A Comparative Study with Commercial Rice Varieties. *Malaysian Applied Biology* 41(1): 29-35.
- Anwar, M. P., Juraimi, A. S., Man, A., Puteh, A., Selamat, A. & Begum, M. (2010). Weed Suppressive Ability of Rice (*Oryza sativa* L.) Germplasm under Aerobic Soil Conditions. *Australian Journal of Crop Science* 4: 706-717.
- Arriaga, L., Huerta, E., Lira-Saade, R., Moreno, E. & Alarcon, J. S. (2006). Assessing the Risk of Releasing Transgenic *Cucurbita* Spp. in Mexico. *Agriculture, Ecosystems and Environment* 112: 291-299.
- Azmi, M., Azlan, S., Yim, K. M., George, T. V. & Chew, S. E. (2012). Control of Weedy Rice in Direct-Seeded Rice Using the Clearfield Production System in Malaysia. *Pak. J. Weed Science Resistance* 18: 49-53.
- Beckie, H. J. & Hall, L. M. (2014). Genetically-Modified Herbicide Resistant (GMHR) Crops a Two Edge Sword? An American Perspective on Development and Effect on Weed Management. *Crop Protection* 66: 40-65.
- Burgos, N R., Norman, R. J., Gealy, D. R. & Black, H. (2006). Competitive N Uptake between Rice and Weedy Rice. *Field Crop Research* 99: 96-105.
- Burgos, N. R., Shivrain V. K., Scott R. C., Mauromoustakos A., Kuk Y.I., Sales M. A. & Bullington J. (2011). Differential Tolerance of Weedy Red Rice (*Oryza sativa* L.) from Arkansas, USA to Glyphosate. *Crop Protection* 30: 986-994.
- Busconi, M., Rossi, D., Lorenzoni, G., Baldi, G. & Fogher, C. (2012). Spread of Herbicide-Resistant Weedy Rice (red rice, *Oryza sativa* L.) after 5 Years of Clearfield Rice Cultivation in Italy. *Plant Biology* 14(5): 751-759.
- Camargo, E. R., Senseman, S. A., McCauley, G. N. & Guice, J. B. (2011). Rice Tolerance to Saflufenacil in Clomazone Weed Control Program. *International Journal of Agronomy*. Doi:10.1155/2011/402461.
- Celec, P., Kukučková, M., Renczésóvá, V., Natarajan S., Pálffy R., Gardlík R., Hodosy J., Behuliak M., Vlková B., Minárik G., Szemes T., Stuchlík S. & Turňa J. (2005). Biological and Biomedical Aspects of Genetically Modified Food. *Biomedicine & Pharmacotherapy* 59: 531-540.
- Chin D.V. (2001). Biology and Management of Barnyard Grass, Red Sprangletop and Weedy Rice. *Weed Biology Management* 1: 37-41.
- Chin, D. V., Thien T. C., Bi H. H. & Nhiem N. T. (2007). Study on Weed and Weedy Rice Control by Imidazolinone Herbicides in Clearfield™ Paddy Grown by Imi- Tolerance Indica Rice Variety. *Omonrice* 15: 63-67.

- Colbach, N., Devauxd, C. & Angevine, F. (2009). Comparative Study of the Efficiency of Buffer Zones and Harvest Discarding on Gene Flow Containment in Oilseed Rape. A Modelling Approach. *European Journal of Agronomy* 30: 187-198.
- Croughan, T. P. (2003). Clearfield Rice: It's Not a GMO. *Louisiana Agriculture Magazine*, Fall Issue.
- Conner, A. J. & Jacobs, J. M. E. (1999). Genetic Engineering of Crops as Potential Source of Genetic Hazard in the Human Diet. *Mutation Research* 443: 223-234.
- Demont, M. & Stein, A. J. (2013). Global Value of GM Rice: A Review of Expected Agronomic and Consumer Benefits. *New Biotechnology* 30(5): 426-436.
- FAO, (1999). *Report of the Global workshop on red rice control*. Varadero, Cuba, 30 August-3 September, pp. 55.
- FAO, (2014a). *FAO Estimates 2014-2015 Global Rice Production to Decline Slightly y/y to 495.6 Million Tons*. Retrieved February 20, 2015, from <http://www.oryza.com/news/usda-lowers-2013-14-global-rice-production-forecast-36-million-tons>.
- FAO, (2014b). *FAO Estimates 2014 China Paddy Rice Production to Increase to 207.44 Million Tons, Up 2% from Last Year*. Retrieved February 20, 2015 from <http://www.oryza.com/news/rice-news/fao-estimates-2014-china-paddy-rice-production-increase-20744-million-tons-2-last>.
- FAO, (2014c). *FAO Estimates India 2014 Paddy Rice Production at 157.7 Million Tons, Down 1% from Last Year*. Retrieved February 20, 2015, from <http://www.oryza.com/news/rice-news/fao-estimates-india-2014-paddy-rice-production-1575-million-tons-down-1-last-year>.
- FAO, (2014d). *FAO Estimates Indonesia 2014 Imports at 1.1 Million Tons; Up 60% from Last Year*. Retrieved February 20, 2015 from <http://www.oryza.com/news/rice-news/fao-estimates-indonesia-2014-imports-11-million-tons-60-last-year>.
- FAO, (2014e). *FAO Estimates Vietnam Rice Exports Will Increase to 7.2 in 2014, up 8% from Last Year*. Retrieved February 20, 2015 from <http://www.oryza.com/news/rice-news/fao-estimates-vietnam-rice-exports-will-increase-72-million-tons-2014-8-previous-year>.
- FAO, (2014f). *FAO Forecasts Thailand 2015 Rice Exports at 11 Million Tons, Up 7% from Last Year*. Retrieved February 20, 2015 from <http://www.oryza.com/news/rice-news-asia-pacific/fao-forecasts-thailand-2014-rice-exports-96-million-tons-45-last-year>.

- FAO, (2014g). *FAO Estimates Malaysia 2014-15 Rice Imports at 1.1 Million Tons, Up 10% from Last Year*. Retrieved February 20, 2015 from <http://www.oryza.com/news/rice-news/fao-estimates-malaysia-2014-15-rice-imports-11-million-tons-10-last-year>.
- Fory, L., Gonzalez, E., Velasquez, A.M., Morales, M., Arcia, K., Blanco, A. E., Quintero, M., Ortiz, A., Perez, I., Duque, M. C. & Lentini, Z. (2006). Gene Flow Analysis in Rice Wild / Weedy Relatives in Tropical America: Understanding Crop. *Biodiversity Interactions*, International Centre for Tropical Agriculture.
- Fragiorge, E. J., Rezende, A. A. A.D., Graf, U. & Spano, M. A. (2008). Comparative Genotoxicity Evaluation of Imidazolinone Herbicides in Somatic Cells of *Drosophila melanogaster*. *Food and Chemical Toxicology* 46: 393–401.
- Garcia, A. A. F., Benchimol, L. L. B., Barbosa, A. M. M., Geraldi, I. O., Souza, C. L. & Souza, A. P. (2004). Comparison of RAPD, RFLP, AFLP and SSR Markers for Diversity Studies in Tropical Maize Inbred Line. *Genetics and Molecular Biology* 27(4): 579-588.
- Garris, A. J., Tai T. H., Coburn, J., Kresovich, S. & McCouch, S. (2005). Genetic Structure and Diversity in *Oryza sativa* L. *Genetics* 169: 1631–1638.
- Gealy, D. R., Mitten, D. H. & Rutger, J. N. (2003). Gene Flow between Red Rice (*Oryza sativa*) and Herbicide-Resistant Rice (*O. sativa*): Implications for Weed Management. *Weed Technology* 17: 627-645.
- Goulart, I. C. G. D. R., Pacheco, M. T., Nunes, A. L. & Merotto A. J. (2012). Identification of Origin and Analysis of Population Structure of Field-Selected Imidazolinone-Herbicide Resistant Red Rice (*Oryza sativa*). *Euphytica* 187: 437-447.
- Gressel, J. & Valverde, B.E. (2009). A Strategy to Provide Long-Term Control of Weedy Rice while Mitigating Herbicide Resistance Transgene Flow, and its Potential Use for Other Crops with Related Weeds, *Pest Management Science* 65: 723–731.
- Gross, B. L., Reagon, M., Hsu, S. C., Caicedo, A. L., Jia, Y. & Olsen, K. M. (2010). Seeing Red: The Origin of Grain Pigmentation in US Weedy Rice. *Molecular Ecology* 19: 3380-3393.
- Guadagnuolo, R., Savova-Bianchi D. & Felber F. (2001). Gene Flow from Wheat (*Triticum aestivum* L.) to Jointed Goatgrass (*Aegilops cylindrical Host*), as Revealed by RAPD and Microsatellite Markers. *Theoretical and Applied Genetics* 103: 1-8.
- Hamid, Z. A. A., Mansor, M. & Man, A. (2007). Life Cycle and Morphological Characteristics of Weedy Rice (*Oryza sativa*) Complex, Locally Called Padi Angin), a Noxious Weed of Rice fields in Malaysia. *Jurnal Biosains* 18(1): 55-79.

- Hansen, M. K. (1998). *Genetic Engineering is Not an Extension of Conventional Plant Breeding*. Retrieved December 1, 2011, from <http://www.consumersunion.org/food/widecpi200.htm>.
- Hernandez-Campuzano, B., Suarez, R., Lina, L., Hernandez, V., Villegas, E., Corzo, G. & Iturriaga, G. (2009). Expression of a Spider Venom Peptide in Transgenic Tobacco Confers Insect Resistance. *Toxicon* 53: 122-128.
- Huangfu, C., Qiang, S. & Song, X. (2011). Performance of Hybrids between Transgenic Oilseed Rape (*Brassica napus*) and Wild *Brassica juncea*: An Evaluation of Potential for Transgene Escape. *Crop Protection* 30: 57-62.
- Indian GMO research Information system (IGMORIS). (2011). Retrieved November 25, 2014 from [//igmoris.nic.in/Files%5CBiologyDocuments%5CBiology_of_Rice.pdf](http://igmoris.nic.in/Files%5CBiologyDocuments%5CBiology_of_Rice.pdf).
- International Rice Research Institute (IRRI). (2002). Standard evaluation system for rice. Manila: International Rice Research Institute, Rice Knowledge Bank. Retrieved February 10, 2013 from <http://www.knowledgebank.irri.org/extension/index.php/seshtm>.
- International Service for the Acquisition of Agri-Biotech Applications (ISAAA), (2014). Pocket K No. 17: Genetic Engineering and GM Crops. Retrieved February 21, 2015 from www.isaaa.org/resources/publications/pocketk/17/default.asp.
- Ishikawa, R., Toki, N., Imai, K., Sato, Y. I., Yamagishi, H., Shimamoto, Y., Ueno, K., Morishima, H. & Sato T. (2005). Origin of Weedy Rice Grown in Bhutan and the Force of Genetic Diversity. *Genetic Resources and Crop Evolution* 6:395-403.
- Juan, Z., Burgos, N. R., Kun, M., Jun, Z. Y., Mei, G., R. & Qing, Y. L. (2008). Genetic Diversity and Relationship of Weedy Rice in Taizhou City, Jiangsu Province, China. *Rice Science* 15(4): 295-302.
- Kane, N. C. & Baack, E. J. (2007). The Origins of Weedy Rice. *Molecular Ecology*, 21: 4423-4425.
- Kaloumenos, N. S., Capote, N., Aguado, A. & Eleftherohorinos, I. G. (2013). Red Rice (*Oryza sativa*) Cross-Resistance to Imidazolinone Herbicides used in Resistant Rice Cultivars Grown in Northern Greece. *Pesticide Biochemistry and Physiology* 105: 177-183.
- Karim, R. S. M., Man, A. B. & Sahid, I. B. (2004). Weed Problems and Their Management in Rice Fields of Malaysia: An Overview. *Weed Biology and Management* 4: 177-186.
- Kumar, V., Bellinder, R.R., Brainard, D.C., Malik, R.K. & Gupta, R.K. (2008). Risks of Herbicide-Resistant Rice in India: A Review. *Crop Protection* 83: 320-329.

- Kwon, S. L. Smith, R. J. & Talbert, R. H. (1992). Comparative Growth and Development of Red Rice (*Oryza sativa*) and Rice (*O. sativa*). *Weed Science* 40: 57-62.
- Lang, N. T. & Buu B. C. (2007). Rice Breeding and Inheritance of Herbicide Resistance in Clearfield Rice. *Omonrice* 15: 36-45.
- Li, L. Y., Xi, Y. X., Ping, Z. F. & Hui, X. M. (2006). SSR Marker Analysis on Indica-Japonica Differentiation of Natural Population of *Oryza rufipogon* in Yuanjiang, Yunnan Province. *Rice Science* 12: 71-74.
- Liang, L. Y. (2006). Exploration of Biopolitics of GMOs: Using Golden Rice as an Analytical Model. *Agriculture Science in China* 12: 885-894.
- Lu, B. R. & Snow, A. A. (2005). Gene Flow from Genetically Modified Rice and its Environmental Consequences. *BioScience*, 55(8): 669-678.
- Lu, B. R. & Yang, C. (2009). Gene Flow from Genetically Modified Rice to its Wild Relatives: Assessing Potential Ecological Consequences. *Biotechnology Advances* 27: 1083-1091.
- Makerly, H., Rahmat, Z. & Wagiran, A. (2012). Potential Use of Partial Desiccation Treatment for Regeneration System of Malaysian Indica Rice (*O. sativa* L.) *Jurnal Teknologi (Sciences & Engineering)* 59: 97-100.
- Mandaokar, A. D., Goyal, R. K., Shukla, A., Bisaria, S., Bhalla, R., Reddy, V.S., Chaurasia, A., Sharma, R.P., Altosaar, I. & Kumar, P. A. (2000). Transgenic Tomato Plants Resistant to Fruit Borer (*Helicoverpa armigera* Hubner). *Crop Protection* 19: 307-312.
- Marembe, B. (2009). Weedy Rice – Evolution, Threats and Management. *Tropical Agriculturist* 157:43-64.
- Nelson, G. C. (2001). *Genetically Modified Organisms in Agriculture : Economics and Politics* pp 7, 9-13, 26-27, 50-51, 200-201, 287. London, Academic Press.
- Normile, D. (2014). China Pulls Plug on Genetically Modified Rice and Corn. Retrieved December 18, 2014 from <http://news.sciencemag.org/asiapacific/2014/08/china-pulls-plug-genetically-modified-rice-and-corn>.
- Olofsdotter, M., Valverde, B. E. & Madsen, K. H. (2000). Herbicide Resistant Rice (*Oryza sativa* L.): Global Implications for Weedy Rice and Weed Management. *Annals of Applied Biology* 137: 279-295.
- Park, J., McFarlane, I., Phipps, R. & Ceddia, G. (2011). The Impact of the EU Regulatory Constraint of Transgenic Crops on Farm Income. *New Biotechnology* 28: 396-406.

- Pintado, S., Montoya, M. R. & Mellado, J. M. R. G. (2011). Imidazolinone Herbicides in Strongly Acidic Media: Speciation and Electroreduction. *Comptes Rendus Chimie* 14: 957-962.
- Peng, S., Huang, J., Sheehy, J.E., Laza, R.C., Visperas, R.M., Zhong, X., Centeno, G.S., Khush, G.S. & Cassman, K.G. (2004). Rice Yields Decline with Higher Night Temperature from Global Warming. *Proceedings of the National Academy of Sciences USA* 101: 9971-9975.
- Perera, U. I. P., Ratnasekera, W. A. D. P. R. & Senanayake, S. G. J. N. (2010). Morphological Diversity of Weedy Rice Accessions Collected In Ampara District In *15th International Forestry and Environment Symposium*. University of Sri Jayewardenepura, Sri Lanka.
- Rajguru, S. N., Burgos, N. R., Stewart, J. M. & Gealy, D. (2002). Genetic Diversity in Red Rice using SSR Markers. *Proceedings of Weed Science Society of America*. 55:115-116.
- Rajguru, S. N., Burgos, N. R., Shivrain, V. K., Stewart, J. M. (2005) Mutations in the Red Rice ALS Gene Associated with Resistance to Imazethapyr. *Weed Science* 53(5): 567-577
- Ramesh, S., Nagadhara, D., Pasalu, I. C., Kumari, A. P., Sarma, N. P., Reddy, V. D. & Rao, K. V. (2004). Development of Stem Borer Resistant Transgenic Parental Lines Involved in the Production of Hybrid Rice. *Journal of Biotechnology* 111: 131-141.
- Ramezani, M., Simpson, N., Oliver, D., Kookana, R., Gill, G. & Preston, C. (2009). Improved Extraction and Clean-Up of Imidazolinone Herbicides from Soil Solutions using Different Solid-Phase Sorbents. *Journal of Chromatography A* 1216: 5092-5100.
- Rathore M, Singh R, & Kumar B (2013). Weedy rice: An Emerging Threat to Rice Cultivation and Options for its Management, *Current Science* 105: 1067 - 1072
- Reuter, H., Schmidt, G., der, W. S., Middelhoff, U., Pehlke, H. & Breckling, B. (2011). Regional Distribution of Genetically Modified Organisms (GMOs)—Up-Scaling the Dispersal and Persistence Potential of Herbicide Resistant Oilseed Rape (*Brassica napus*). *Ecological Indicator* 11: 989-999.
- Roso, A.C., Merotto, A. J., Delatorre, C.A. & Menezes, V.G. (2010). Regional Scale Distribution of Imidazolinone Herbicide-Resistant Alleles in Red Rice (*Oryza sativa* L.) Determined Through SNP Markers. *Field Crops Research* 119: 175-182.
- Saito, K., Azoma K. & Rodenburg J. (2010). Plant Characteristics Associated with Weed Competitiveness of Rice under Upland and Lowland Conditions in West Africa. *Field Crops Research* 117: 1-8.

- Sánchez-Olguín, E., Arrieta-Espinoza, G. & Espinoza-Esquivel, A.M. (2007). Vegetative and Reproductive Development of Costa Rican Weedy Rice Compared with Commercial Rice (*Oryza sativa*). *Planta Daninha* 1: 13–23.
- Saragih, A. A., Puteh, A. B., Ismail, M. R. & Mondal, M. M. A. (2013). Pollen Quality Traits of Cultivated (*Oryza sativa* L. ssp. indica) and Weedy (*Oryza sativa* var. nivara) Rice to Water Stress at Reproductive Stage. *Australian Journal of Crop Science* 7(8), 1106-1112.
- Shaner, D.L. & Singh, B.K. (1997). *Herbicide Activity: Toxicology, Biochemistry, and Molecular Biology* pp. 69–110. Washington DC, IOS Press.
- Shimono, H., Hasegawa, T., Moriyama, M., Fujimura, S., Nagata, T. (2005) Modeling Spikelet Sterility Induced by Low Temperature in Rice. *Agronomy Journal* 97: 1524-1536
- Shivrain, V. K., Burgos, N. R., Anders, M. M., Rajguru, S. N., Moore, J. & Sales, M. A. (2007). Gene Flow Between Clearfield™ Rice and Red Rice. *Crop Protection* 26: 349-356.
- Shivrain, V. K., Burgos, N. R., Gealy, D. R., Moldenhauer, K.A. K. & Baquireza, C. J. (2008). Maximum Outcrossing Rate and Genetic Compatibility between Red Rice (*Oryza sativa*) Biotypes and Clearfield™ Rice. *Weed Science* 56:807-813.
- Shivrain, V. K., Burgos, N. R., Gealy, D. R., Sales, M. A. & Smith, K. L. (2009). Gene Flow from Weedy Red Rice (*Oryza sativa* L.) to Cultivated Rice and Fitness of Hybrids. *Pest Management Science* 65: 1124-1129.
- Shivrain, V. K., Burgos, N. R., Sales, M. A. & Kuk, Y. I. (2010). Polymorphism in the ALS Gene of Weedy Rice (*Oryza sativa* L.) Accessions with Differential Tolerance to Imazethapyr. *Crop Protection* 29: 336-341.
- Soleri, D., Cleveland, D. A., Glasgow, G., Sweeney, S. H., Cuevas, F. A., Fuentes, M. R. & Humberto R. L. (2008). Testing Assumptions Underlying Economic Research on Transgenic Food Crops for Third World Farmers: Evidence from Cuba, Guatemala and Mexico. *Ecological Economics* 67: 667-682.
- Song, B. K., Chuah, T. S., Tam, S. M. & Olsen, K. M. (2014). Malaysian Weedy Rice Show its True Stripes: Wild *Oryza* and Elite Rice Cultivars Shape Agriculture Weed Evolution in Southeast Asia. *Molecular Ecology* 23: 5003-5017.
- Song, H. Y., Cho, S. J., Lim, H. K., Park, N. J. & Hwang, I. T. (2010). Transformation a Mutant *Monochoria vaginalis* Acetolactate Synthase (ALS) Gene Renders *Arabidopsis thaliana* Resistant to ALS Inhibitors. *Pesticide Biochemistry and Physiology* 97: 223-228.
- Song, Z. P., Lu, B.-R., Zhu, Y. G. & Chen, J. K. (2003). Gene Flow from Cultivated Rice to the Wild Species *Oryza rufipogon* Under Experimental Field Conditions. *New Phytologist* 157: 657-665.

- Stewart, C. N. J., Halfhill, M. D. & Warwick, S. I. (2003). Transgene Introgression from Genetically Modified Crops to Their Wild Relatives. *National Review Genetics* 4: 806-817.
- Su, J. & Wu, R. (2003). Stress-Inducible Synthesis of Proline in Transgenic Rice Confers Faster Growth Under Stress Conditions than that with Constitutive Synthesis. *Plant Science* 166: 941-948.
- Sudianto E., Song B. K., Neik T. X., Nestor E. S., Robert C. S., & Burgos N. R. (2013) Clearfield® rice: Its Development, Success and Key Challenges on a Global Perspective. *Crop Protection*. 49: 40-51.
- Sweeney, M. & McCouch, S. (2007). The Complex History of the Domestication of Rice. *Annals of Botany* 100: 951-957.
- Thompson, C. J., Thompson, B. J. P., Ades, P. K., Cousens, R., Garnier-Gere, P., Landman, K., Newbigin, E. & Burgman, M. A. (2003). Model-Based Analysis of the Likelihood of Gene Introgression from Genetically Modified Crops into Wild Relatives. *Ecological Modelling* 162: 199-209.
- Vaughan D. A., Morishima H. & Kadowaki K. (2003). Diversity in the *Oryza* Genus. *Current Opinion in Plant Biology* 6:139-146.
- Wang, G. X., Tan, M. K., Rakshit, S., Saitoh, H., Terauchi, R., Imaizumi, T., Ohsako, T. & Tominaga, T. (2007). Discovery of Single-Nucleotide Mutations in Acetolactate Synthase Genes by Ecotilling. *Pesticide Biochemistry and Physiology* 88: 143–148.
- Wang, Y.M., Zhang, G.A., Du, J.P., Liu, B. & Wang, M.C. (2010) Influence of Transgenic Hybrid Rice Expressing a Fused Gene Derived from CryIAb and CryIac on Primary Insect Pests and Rice Yield. *Crop Protection* 29: 128–133.
- Yang, P., Iles, M., Yan, S. & Jolliffe, F. (2005). Farmers' Knowledge, Perceptions and Practices in Transgenic Bt Cotton in Small Producer Systems in Northern China. *Crop Protection* 24: 229-239.
- Yong, I. K., Burgos, N. R., & Shivrain V. K. (2008). Natural Tolerance to Imazethapyr in Red Rice (*Oryza sativa*) *Weed Science* 56:1-11
- Yu, H., Xue-liang, X., Wei-hua, M., Ben-qi, Y., Hui, W., Fang-zhou, L., Man-qun, W., Gang, W. & Hong-xia, H. (2011). The Influence of Transgenic cryIAb/cryIAc, cryIc and cry2A Rice on Non-Target Planthoppers and Their Main Predators under Field Conditions. *Agriculture Sciences in China* 11: 1739-1747.
- Zhang, W. (2005). Risk Assessment of the Transfer of Imazethapyr Herbicide Resistance from Clearfield Rice to Red Rice. In *Department of Agronomy and Environmental Management*, pp. 136: Louisiana State University.